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US	60/172,247 (CIP)
Filed on	11 January 1999 (11.01.99)
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(54) Title: HUMAN PEPTIDASES

(57) Abstract

The invention provides human peptidases (HPEP) and polynucleotides which identify and encode HPEP. The invention also provides expression vectors, host cells, antibodies, agonists, and antagonists. The invention also provides methods for diagnosing, treating, or preventing disorders associated with expression of HPEP.

Table 2 (cont.)

SEQ ID NO:	Amino Acid Residues	Potential Phosphorylation Sites	Potential Glycosylation Sites	Signature Sequences, Motifs, and Domains	Homologous Sequences	Analytical Methods
14	703	S20 S68 T120 T135 S331 T383 S562 S606 S607 S631 S674 S698 T31 S95 S115 S173 S355 S490 S562 S650	N318 N434 N445 N670	E1 ubiquitin activating enzyme: K352-H442	E1-like protein (ubiquitin activating enzyme) [Pichia pastoris] g4262402	MOTIFS BLAST BLIMPS
15	145	T36 S100 S115 T47	N34	Protease serine hydrolase precursor signal zymogen glycoprotein multigene family: L16-Q64, G87-K140 Trypsin: L25-Q64, S84-N142	Matriptase (serine protease) [Homo sapiens] g5359675, g6002714 Epithin (membrane bound serine protease) [Mus musculus] g4104970	MOTIFS BLAST BLIMPS
16	518	S74 T252 S151 T169 T245 S312 S361 T419 S462 S502 S16 S70 S98 S133 T301 S331 S428 T516 Y334	N234	Dipeptidyl peptidase IV: H255-L305, E326-Q352, E379-P411	Dipeptidyl peptidase IV [Stenotrophomonas maltophilia] g1753197	MOTIFS BLAST BLIMPS

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                650                 655                 660
Gly Phe Asn Phe Leu Ala Lys Val Phe Asn Ser Ser His Ser Phe
                665                 670                 675
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                35                  40                  45
Ile Thr Pro Arg Met Met Cys Val Gly Phe Leu Ser Gly Gly Val
                50                  55                  60
Asp Ser Cys Gln Val Ala Pro Gly Ala Gly Gly Arg Gln Val Gly
                65                  70                  75
Pro Gly Arg Gly Gly Thr Gly Asp Ser Pro Ala Gly Leu Val Ser
                80                  85                  90
Ala Gln Gly Asp Ser Gly Gly Pro Leu Ser Ser Val Glu Ala Asp
                95                 100                 105
Gly Arg Ile Phe Gln Ala Gly Val Val Ser Trp Gly Asp Gly Cys
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             50                  55                  60
Asp Gly Leu Arg Ser Ile Ile His Gly Ser Arg Lys Tyr Ser Gly
             65                  70                  75
Leu Ile Val Asn Lys Ala Pro His Asp Phe Gln Phe Val Gln Lys
             80                  85                  90
Thr Asp Glu Ser Gly Pro His Ser His Arg Leu Tyr Tyr Leu Gly
             95                 100                 105
Met Pro Tyr Gly Ser Arg Glu Asn Ser Leu Leu Tyr Ser Glu Ile
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            125                 130                 135
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Ser Arg Glu Glu Glu Leu Leu Arg Glu Arg Lys Arg Leu Gly Val
            155                 160                 165
Phe Gly Ile Thr Ser Tyr Asp Phe His Ser Glu Ser Gly Leu Phe
            170                 175                 180
Leu Phe Gln Ala Ser Asn Ser Leu Phe His Cys Arg Asp Gly Gly
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Lys Asn Gly Phe Met Val Ser Pro Met Lys Pro Leu Glu Ile Lys
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            215                 220                 225
Asp Pro Asp Phe Phe Ser Phe Ile Asn Asn Ser Asp Leu Trp Val
            230                 235                 240
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Gln Gly Leu Ser Asn Val Leu Asp Asp Pro Lys Ser Ala Gly Val
            260                 265                 270
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Trp Trp Cys Pro Thr Ala Ser Trp Glu Gly Ser Glu Gly Leu Lys
            290                 295                 300
Thr Leu Arg Ile Leu Tyr Glu Glu Val Asp Glu Ser Glu Val Glu
            305                 310                 315
Val Ile His Val Pro Ser Pro Ala Leu Glu Glu Arg Lys Thr Asp
            320                 325                 330
Ser Tyr Arg Tyr Pro Arg Thr Gly Ser Lys Asn Pro Lys Ile Ala
            335                 340                 345
Leu Lys Leu Ala Glu Phe Gln Thr Asp Ser Gln Gly Lys Ile Val
            350                 355                 360
Ser Thr Gln Glu Lys Glu Leu Val Gln Pro Phe Ser Ser Leu Phe
            365                 370                 375
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Val Gln Pro Tyr Val Val Tyr Glu Glu Val Thr Asn Val Trp Ile
                440                 445                 450
Asn Val His Asp Ile Phe Tyr Pro Phe Pro Gln Ser Glu Gly Glu
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Asp Glu Leu Cys Phe Leu Arg Ala Asn Glu Cys Lys Thr Gly Phe
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Cys His Leu Tyr Lys Val Thr Ala Val Leu Lys Ser Gln Gly Tyr
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                35                  40                  45
Arg Phe Lys Arg Ala Ile Phe Gln Gly Gln Tyr Cys Arg Asn Phe
                50                  55                  60
Gly Cys Cys Glu Asp Arg Asp Asp Gly Cys Val Thr Glu Phe Tyr
                65                  70                  75
Ala Ala Asn Ala Leu Cys Tyr Cys Asp Lys Phe Cys Asp Arg Glu
                80                  85                  90
Asn Ser Asp Cys Cys Pro Asp Tyr Lys Ser Phe Cys Arg Glu Glu
                95                 100                 105
Lys Glu Trp Pro Pro His Thr Gln Pro Trp Tyr Pro Glu Gly Cys
               110                 115                 120
Phe Lys Asp Gly Gln His Tyr Glu Glu Gly Ser Val Ile Lys Glu
               125                 130                 135
Asn Cys Asn Ser Cys Thr Cys Ser Gly Gln Gln Trp Lys Cys Ser
               140                 145                 150
Gln His Val Cys Leu Val Arg Ser Glu Leu Ile Glu Gln Val Asn
               155                 160                 165
Lys Gly Asp Tyr Gly Trp Thr Ala Gln Asn Tyr Ser Gln Phe Trp
               170                 175                 180
Gly Met Thr Leu Glu Asp Gly Phe Lys Phe Arg Leu Gly Thr Leu
               185                 190                 195
Pro Pro Ser Pro Met Leu Leu Ser Met Asn Glu Met Thr Ala Ser
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